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OM protein - protein search, using sw model

Run on: June 5, 2003, 14:00:01 ; Search time 8.72727 Seconds
(without alignments)
445.022 Million cell updates/sec

Title: Perfect score:
Sequence: US-09-840-795-15
Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 262574 seqs, 29422922 residues

Total number of hits satisfying chosen parameters: 262574

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Maximum Match 0%
Maximum Match 100%
Listing first: 45 summaries

Database : Issued Patents_AA.*
1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*5: /cgn2_6/ptodata/1/iaa/PCNRS_COMB.pep:*6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match Length	DB ID	Description
1	198.5	28.0	210	4 US-09-286-529-3
2	188	26.5	151	4 US-09-286-529-3
3	117	16.5	448	4 US-09-342-681C-17
4	117	16.5	448	4 US-09-342-681C-19
5	96	13.5	155	4 US-09-146-950-4
6	96	13.5	159	4 US-09-146-950-20
7	96	13.5	193	4 US-09-146-950-2
8	96	13.5	197	4 US-09-146-950-18
9	96	13.5	283	4 US-09-509-024-2
10	96	13.5	283	4 US-09-333-279-2
11	96	13.5	283	4 US-09-572-939C-2
12	96	13.5	283	5 PCT-US56-12374-2
13	96	419	4 US-08-509-024-7	
14	96	419	4 US-09-509-024-7	
15	95.5	13.5	121	4 US-08-28-603A-18
16	95.5	13.5	260	4 US-09-006-353A-8
17	89	12.6	260	4 US-09-573-986-8
18	89	12.6	970	2 US-08-173-789-7
19	89	12.6	973	1 US-08-162-809-10
20	89	12.6	988	1 US-08-162-809-14
21	88	12.4	169	3 US-08-330-112-11
22	88	12.4	169	4 US-09-375-119-11
23	87	12.3	176	1 US-08-114-936A-18
24	87	12.3	176	2 US-08-926-922-18
25	87	12.3	176	3 US-09-253-602-18
26	87	12.3	176	4 US-09-527-18
27	86.5	12.2	186	1 US-08-189-438B-6

ALIGNMENTS

```

RESULT 1
US-09-286-529-3
; Sequence 3, Application US/09286529
; Patent No. 6297367

GENERAL INFORMATION:
APPLICANT: Catherine Tribouley
TITLE OF INVENTION: NEW MEMBERS OF TNF AND TNFR FAMILIES
FILE REFERENCE: 1408 003/200130.439C1
CURRENT APPLICATION NUMBER: US/09-286,529
CURRENT FILING DATE: 1999-04-05
NUMBER OF SEQ ID NOS: 25
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO: 3
LENGTH: 210
; TYPE: PRT
; ORGANISM: human
US-09-286-529-3

```

Query Match Best Local Similarity 28.0%; Score 198.5; DB 4; Length 210;
Matches 40; Conservativeness 40.4%; Pred. No. 2.4e-15; Indels 9; Gaps 2;

QY 2 DCQENFYWDQGRGRCVTCQRCCPGQELSKDCCGSGEGGDAYCAGCPPRSTKAGATNNRVA 61
Db 33 DCRQQEFKDRSGNCVLLCKQCGPGMELSKCECGFGYGEDAQCVCPRHFKEDMGFQKCP 92
QY 62 SPVLSSIVVERFRNFTMSXAVKQ---GXFXQVSNRTR 96
Db 93 ADCALVNRFQORANCSHTSDAVGDCLPGFY----RKT 126

RESULT 2
US-09-286-529-4
; Sequence 4, Application US/09286529
; Patent No. 6297367

GENERAL INFORMATION:
APPLICANT: Catherine Tribouley
TITLE OF INVENTION: NEW MEMBERS OF TNF AND TNFR FAMILIES
FILE REFERENCE: 1408 003/200130.439C1
CURRENT APPLICATION NUMBER: US/09-286,529
CURRENT FILING DATE: 1999-04-05
NUMBER OF SEQ ID NOS: 25
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO: 4
LENGTH: 151
; TYPE: PRT
; ORGANISM: human
US-09-286-529-4

Query Match 26.5%; Score 188; DB 4; Length 151;

Best Local Similarity 40.0%; Pred. No. 2.7e-14; Mismatches 36; Indels 10; Gaps 3; Matches 40; Conservative 14; Mismatches 36; Indels 10; Gaps 3;

RESULT 3
US-09-342-681C-17
; Sequence 17, Application US/09342681C
; Patent No. 635582
; GENERAL INFORMATION:
; APPLICANT: Zonana et al.
; TITLE OF INVENTION: Hypohidrotic ectodermal dysplasia genes and proteins
FILE REFERENCE: 52978
CURRENT APPLICATION NUMBER: US/09/342,681C
CURRENT FILING DATE: 1999-06-29
PRIOR APPLICATION NUMBER: 60/092,279
PRIOR FILING DATE: 1998-07-09
PRIOR APPLICATION NUMBER: 60/112,366
PRIOR FILING DATE: 1998-12-15
NUMBER OF SEQ ID NOS: 123
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 17
LENGTH: 448
TYPE: PRT
ORGANISM: Homo sapiens
US-09-342-681C-17

Query Match 16.5%; Score 117; DB 4; Length 448;
Best Local Similarity 42.6%; Pred. No. 1.6e-05; Mismatches 23; Conservative 5; Indels 24; Gaps 2; Matches 23; Conservative 5; Mismatches 24; Indels 2; Gaps 2;

QY 2 DCQENBYWDO-WGRCVTCQRCGPQGQELSKDGYGGDDAY-CTACPPRSTKAAG 53
Db 30 NCGENEYNYNQTTGLCQCPGPGPEPYLSCGGYGTKEAEGELTG--TVCEPCPPGTYIAHNGLSKLOC 83

RESULT 4
US-09-342-681C-19
; Sequence 19, Application US/09342681C
; Patent No. 635582
; GENERAL INFORMATION:
; APPLICANT: Zonana et al.
; TITLE OF INVENTION: Hypohidrotic ectodermal dysplasia genes and proteins
FILE REFERENCE: 52978
CURRENT APPLICATION NUMBER: US/09/342,681C
CURRENT FILING DATE: 1999-06-29
PRIOR APPLICATION NUMBER: 60/092,279
PRIOR FILING DATE: 1998-07-09
PRIOR APPLICATION NUMBER: 60/112,366
PRIOR FILING DATE: 1998-12-15
NUMBER OF SEQ ID NOS: 123
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 19
LENGTH: 448
TYPE: PRT
ORGANISM: Homo sapiens
US-09-342-681C-19

Query Match 16.5%; Score 117; DB 4; Length 448;
Best Local Similarity 42.6%; Pred. No. 1.6e-05; Mismatches 23; Conservative 5; Mismatches 24; Indels 2; Gaps 2;

QY 2 DCQENBYWDO-WGRCVTCQRCGPQGQELSKDGYGGDDAY-CTACPPRSTKAAG 53
Db 30 NCGENEYNYNQTTGLCQCPGPGPEPYLSCGGYGTKEAEGELTG--TVCEPCPPGTYIAHNGLSKLOC 83

RESULT 5
US-09-146-950-4
; Sequence 4, Application US/09146950A
; Patent No. 6287808
; GENERAL INFORMATION:
; APPLICANT: Busfield, Samantha J.
; TITLE OF INVENTION: PROTEIN FAMILY AND USES THEREOF
FILE REFERENCE: 0904/057001
CURRENT APPLICATION NUMBER: US/09/146,950A
CURRENT FILING DATE: 1998-09-03
NUMBER OF SEQ ID NOS: 25
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 20
LENGTH: 159
TYPE: PRT
ORGANISM: Homo sapiens
US-09-146-950-4

Query Match 13.5%; Score 96; DB 4; Length 155;
Best Local Similarity 28.9%; Pred. No. 0.0012; Mismatches 28; Conservative 11; Indels 14; Gaps 6; Matches 28; Conservative 11; Mismatches 44; Indels 14; Gaps 6;

QY 3 CQENYWDQWGRCVTCQRCGPQGQELSKDGYGGDDAYCTACPPRS---TAAAGATTNV 58
Db 4 CKEDBY-PVSEC--CPKCSPGYRKAEAGELTG--TVCEPCPPGTYIAHNGLSKLOC 58

QY 59 RVASVLSSIVFRRNCTXAVCG--GXFAQWSN 92
Db 59 QMCDAAMGLRASR-NCSRTPENAVGCCSPGHFCITVQD 93

RESULT 6
US-09-146-950-20
; Sequence 20, Application US/09146950A
; Patent No. 6287808
; GENERAL INFORMATION:
; APPLICANT: Busfield, Samantha J.
; TITLE OF INVENTION: NOVEL MOLECULES OF THE HERPESVIRUS-ENTRY-MEDIATOR-RELATED FILE REFERENCE: 0904/057001
CURRENT APPLICATION NUMBER: US/09/146,950A
CURRENT FILING DATE: 1998-09-03
NUMBER OF SEQ ID NOS: 25
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 20
LENGTH: 159
TYPE: PRT
ORGANISM: Homo sapiens
US-09-146-950-20

Query Match 13.5%; Score 96; DB 4; Length 159;
Best Local Similarity 28.9%; Pred. No. 0.0012; Mismatches 28; Conservative 11; Indels 14; Gaps 6; Matches 28; Conservative 11; Mismatches 44; Indels 14; Gaps 6;

QY 3 CQENYWDQWGRCVTCQRCGPQGQELSKDGYGGDDAYCTACPPRS---TAAAGATTNV 58
Db 4 CKEDBY-PVSEC--CPKCSPGYRKAEAGELTG--TVCEPCPPGTYIAHNGLSKLOC 58

QY 59 RVASVLSSIVFRRNCTXAVCG--GXFAQWSN 92
Db 59 QMCDAAMGLRASR-NCSRTPENAVGCCSPGHFCITVQD 93

RESULT 7
US-09-146-950-2
; Sequence 2, Application US/09146950A
; Patent No. 6287808
; GENERAL INFORMATION:
; APPLICANT: Busfield, Samantha J.
; TITLE OF INVENTION: NOVEL MOLECULES OF THE HERPESVIRUS-ENTRY-MEDIATOR-RELATED FILE REFERENCE: 0904/057001
TITLE OF INVENTION: PROTEIN FAMILY AND USES THEREOF

FILE REFERENCE: 09404/057001
 CURRENT FILING DATE: 1998-09-03
 CURRENT NUMBER: US/09/146,950A
 NUMBER OF SEQ ID NOS: 25
 SOFTWARE: FastSEQ for Windows Version 3.0
 SEQ ID NO 2
 LENGTH: 193
 TYPE: PRT
 ORGANISM: Homo sapiens
 US-09-146-950-2

Query Match 13.5%; Score 96; DB 4; Length 193;
 Best Local Similarity 28.9%; Pred. No. 0.0015; Indels 14; Gaps 6;
 Matches 28; Conservative 11; Mismatches 44; Indexes 44; Gaps 6;

Best Local Similarity 28.9%; Pred. No. 0.0015; Indels 14; Gaps 6;
 Matches 28; Conservative 11; Mismatches 44; Indexes 44; Gaps 6;

Qy 3 COENEWFQDGRCVTCQRCGPQEQELSKDCGEGGDAYCTACPPRS---TKAAGATNV 58
 Db 42 CKEDEV-PVGSEC--CPKCSGPGRYKEACGELTG--TVCBPCPGTYIAHNGLSKCLQC 96

Qy 59 RVASPVLSSTVFRRENCTXTSXAVG--GXFAQVSN 92
 Db 97 QMCDPAMGLRASR-NCSRTENAVCCGSPGHFCIVQD 131

RESULT 8
 US-09-146-950-18
 Sequence 18, Application US/09146950A
 Patent No. 6287808
 GENERAL INFORMATION:
 APPLICANT: Busfield, Samantha J.
 TITLE OF INVENTION: NOVEL MOLECULES OF THE HERPESVIRUS-ENTRY-MEDIATOR-RELATED
 TITLE OF INVENTION: PROTEIN FAMILY AND USES THEREOF
 FILE REFERENCE: 09404/057001
 CURRENT APPLICATION NUMBER: US/09/146,950A
 CURRENT FILING DATE: 1998-09-03
 NUMBER OF SEQ ID NOS: 25
 SOFTWARE: FastSEQ for Windows Version 3.0
 SEQ ID NO 18
 LENGTH: 197
 TYPE: PRT
 ORGANISM: Homo sapiens
 US-09-146-950-18

Query Match 13.5%; Score 96; DB 4; Length 197;
 Best Local Similarity 28.9%; Pred. No. 0.0016; Indels 14; Gaps 6;
 Matches 28; Conservative 11; Mismatches 44; Indexes 44; Gaps 6;

Qy 3 COENEWFQDGRCVTCQRCGPQEQELSKDCGEGGDAYCTACPPRS---TKAAGATNV 58
 Db 42 CKEDEV-PVGSEC--CPKCSGPGRYKEACGELTG--TVCBPCPGTYIAHNGLSKCLQC 96

Qy 59 RVASPVLSSTVFRRENCTXTSXAVG--GXFAQVSN 92
 Db 97 QMCDPAMGLRASR-NCSRTENAVCCGSPGHFCIVQD 131

RESULT 9
 US-09-509-024-2
 Sequence 2, Application US/08509024B
 Patent No. 6291207
 GENERAL INFORMATION:
 APPLICANT: SPEAR, Patricia G.
 APPLICANT: MONTGOMERY, Rebecca I.
 TITLE OF INVENTION: HERPES VIRUS ENTRY RECEPTOR PROTEIN
 FILE REFERENCE: 0290-1
 CURRENT APPLICATION NUMBER: US/08/509,024B
 CURRENT FILING DATE: 1995-07-25
 NUMBER OF SEQ ID NOS: 7
 SOFTWARE: PatentIn Ver. 2.0
 SEQ ID NO 2
 LENGTH: 283
 TYPE: PRT
 ORGANISM: Homo sapiens

RESULT 10
 US-09-333-279-2
 Sequence 2, Application US/09333279
 Patent No. 6303336
 GENERAL INFORMATION:
 APPLICANT: SPEAR, Patricia G.
 APPLICANT: MONTGOMERY, Rebecca I.
 TITLE OF INVENTION: HERPES VIRUS ENTRY RECEPTOR PROTEIN
 FILE REFERENCE: 0290-1
 CURRENT APPLICATION NUMBER: US/09/333,279
 NUMBER OF SEQ ID NOS: 7
 SOFTWARE: PatentIn Ver. 2.0
 SEQ ID NO 2
 LENGTH: 283
 TYPE: PRT
 ORGANISM: Homo sapiens
 US-09-333-279-2

Query Match 13.5%; Score 96; DB 4; Length 283;
 Best Local Similarity 28.9%; Pred. No. 0.0024; Indels 14; Gaps 6;
 Matches 28; Conservative 11; Mismatches 44; Indexes 44; Gaps 6;

Qy 3 COENEWFQDGRCVTCQRCGPQEQELSKDCGEGGDAYCTACPPRS---TKAAGATNV 58
 Db 42 CKEDEV-PVGSEC--CPKCSGPGRYKEACGELTG--TVCBPCPGTYIAHNGLSKCLQC 96

Qy 59 RVASPVLSSTVFRRENCTXTSXAVG--GXFAQVSN 92
 Db 97 QMCDPAMGLRASR-NCSRTENAVCCGSPGHFCIVQD 131

RESULT 11
 US-09-072-993C-2
 Sequence 2, Application US/09072993C
 Patent No. 6346388
 GENERAL INFORMATION:
 APPLICANT: Michael R. Brigham-Burke
 APPLICANT: Peter R. Young
 TITLE OF INVENTION: A METHOD OF IDENTIFYING AGONIST AND
 TITLE OF INVENTION: ANTAGONISTS FOR TUMOR NECROSIS RELATED RECEPTORS TR1 AND TR2
 FILE REFERENCE: GH-50030
 CURRENT APPLICATION NUMBER: US/09/072,993C
 CURRENT FILING DATE: 1998-05-06
 PRIOR APPLICATION NUMBER: 60/055,513
 PRIOR FILING DATE: 1997-08-13
 PRIOR APPLICATION NUMBER: 60/056,980
 PRIOR FILING DATE: 1997-08-26
 PRIOR APPLICATION NUMBER: 60/057,550
 PRIOR FILING DATE: 1997-08-29
 NUMBER OF SEQ ID NOS: 9
 SOFTWARE: FastSEQ for Windows Version 3.0
 SEQ ID NO 2
 LENGTH: 283
 TYPE: PRT
 ORGANISM: HOMO SAPIENS
 US-09-072-993C-2

Query Match 13.5%; Score 96; DB 4; Length 283;
 Best Local Similarity 28.9%; Pred. No. 0.0024; Mismatches 28; Conservative 11; Indels 14; Gaps 6; Matches 28; Mismatches 44; Number of SEQ ID NOS: 7

Y 3 COENEYWQWGRGCYTCQCGPGQBLSKCGYGGDAYCTACPRS ---TKAGATTNV 58
 b 42 CKEDEY-PVGSEC---CPKCSPGYRKEAGELTG---TVCPCPGTYIAHNLGSKCLQC 96
 Y 59 RVASPVLLSIVFRFNCXTSXAVCG--GXFAQVSN 92
 b 97 QMCDPAMGLRASR--NCSRTENAVCGCSPGHFCIVQD 131

RESULT 12
 CT-US96-12374-2
 Sequence 2, Application PC/TUS9612374
 GENERAL INFORMATION:
 APPLICANT: Northwestern University
 TITLE OF INVENTION: Herpes Virus Entry Mediator
 NUMBER OF SEQUENCES: 5
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: Dressler, Goldsmith, Minamow & Katz, Ltd.
 STREET: 180 N. Stetson, Suite 4700
 CITY: Chicago
 STATE: Illinois
 COUNTRY: U.S.A.
 ZIP: 60601

COMPUTER READABLE FORM:
 MEDIUM TYPE: FLOPPY DISK
 COMPUTER: IBM PC COMPATIBLE
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.30
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: PCT/US96/12374
 FILING DATE:
 CLASSIFICATION:

ATTORNEY/AGENT INFORMATION:
 NAME: Northrup, Thomas E.
 REGISTRATION NUMBER: 33,268
 REFERENCE/DOCKET NUMBER: NOR3446P020PC
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: (312) 616-5400
 TELEFAX: (312) 616-5460

INFORMATION FOR SEQ ID NO: 2:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 283 amino acids
 TYPE: amino acid
 TOPOLOGY: linear
 MOLECULE TYPE: protein
 CP-T-US96-12374-2

Query Match 13.5%; Score 96; DB 4; Length 419;
 Best Local Similarity 28.9%; Pred. No. 0.0039; Mismatches 28; Conservative 11; Indels 14; Gaps 6; Matches 28; Mismatches 44; Number of SEQ ID NOS: 7

Y 3 COENEYWQWGRGCYTCQCGPGQBLSKCGYGGDAYCTACPRS ---TKAGATTNV 58
 b 42 CKEDEY-PVGSEC---CPKCSPGYRKEAGELTG---TVCPCPGTYIAHNLGSKCLQC 96
 Y 59 RVASPVLLSIVFRFNCXTSXAVCG--GXFAQVSN 92
 b 97 QMCDPAMGLRASR--NCSRTENAVCGCSPGHFCIVQD 131

RESULT 14
 US-09-333-279-7
 Sequence 7, Application US/09333279
 PATENT NO. 6303336
 GENERAL INFORMATION:
 APPLICANT: SPEAR, Patricia G.
 APPLICANT: MONTGOMERY, Rebecca I.
 TITLE OF INVENTION: HERPES VIRUS ENTRY RECEPTOR PROTEIN
 FILE REFERENCE: 090-1
 CURRENT APPLICATION NUMBER: US/09/333,279
 CURRENT FILING DATE: 1999-06-15
 NUMBER OF SEQ ID NOS: 7
 SOFTWARE: PatentIn Ver. 2.0
 SEQ ID NO. 7
 LENGTH: 419
 TYPE: PRT
 ORGANISM: Homo sapiens
 US-09-333-279-7

Query Match 13.5%; Score 96; DB 4; Length 419;
 Best Local Similarity 28.9%; Pred. No. 0.0039; Mismatches 28; Conservative 11; Indels 14; Gaps 6; Matches 28; Mismatches 44; Number of SEQ ID NOS: 7

Y 3 COENEYWQWGRGCYTCQCGPGQBLSKCGYGGDAYCTACPRS ---TKAGATTNV 58
 b 42 CKEDEY-PVGSEC---CPKCSPGYRKEAGELTG---TVCPCPGTYIAHNLGSKCLQC 96
 Y 59 RVASPVLLSIVFRFNCXTSXAVCG--GXFAQVSN 92
 b 97 QMCDPAMGLRASR--NCSRTENAVCGCSPGHFCIVQD 131

RESULT 15
 US-08-838-683A-18
 Sequence 18, Application US/08828683A
 PATENT NO. 6469144
 GENERAL INFORMATION:
 APPLICANT: Ashkenazi, Avi J.
 TITLE OF INVENTION: APO-2 LI AND APO-3 POLYPEPTIDES
 NUMBER OF SEQUENCES: 28
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: Genentech, Inc.
 STREET: 1 DNA WAY
 CITY: South San Francisco
 STATE: California
 COUNTRY: USA
 ZIP: 94080

COMPUTER READABLE FORM:
 MEDIUM TYPE: 3.5 inch, 1.44 Mb floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Winpatin (Genentech)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/828,683A
FILING DATE: 31-Mar-1997
CLASSIFICATION: <Unknown>

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/625328
FILING DATE: 1-Apr-1996
APPLICATION NUMBER: 08/710802
FILING DATE: 23-Sep-1996
ATTORNEY/AGENT INFORMATION:
NAME: Marschang, Diane L.
REGISTRATION NUMBER: 35,600
REFERENCE/DOCKET NUMBER: P1007P1

TELECOMMUNICATION INFORMATION:

TELEPHONE: 650/225-5416
TELEFAX: 650/952-9881

INFORMATION FOR SEQ ID NO: 18:

SEQUENCE CHARACTERISTICS:

LENGTH: 121 amino acids

TYPE: Amino Acid

TOPOLOGY: Linear

SEQUENCE DESCRIPTION: SEQ ID NO: 18:

US-08-828-683A-18
Query Match 13.5%; Score 95.5; DB 4; Length 121;

Best Local Similarity 34.9%; Pred. No. 0.001; Matches 29; Conservative 5; Mismatches 42; Indels 7; Gaps 4;

Matches 3 COENEYWMDQWGRCVTCORCGPGQELSKDCGGYEGGDAYCTACPPRSTKAAATTNVRAS 62
Db 2 CPERHWTWQKLCL--COPCEPGTFLYKDCDQRKA-RQCDPCIPGVFSFSPDHTRPHCES 58

Qy 63 --PVLSIVVERFRFNCTTISXAVC 83
Db 59 CRHCNSGLILVR--NCTITANAE 79

Search completed: June 5, 2003, 14:00:34
Job time : 9.7227 secs

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